

#11

Variable	Mean	SD	Median	Mode	Range	Skewness	Kurtosis	Shapiro-Wilk	Levene
Age	35.2	12.5	32.0	30.0	20-55	0.15	2.10	0.98	0.95
Gender	1.2	0.4	1.0	1.0	1-2	0.05	0.10	0.99	0.98
Marital Status	2.5	1.0	2.0	2.0	1-4	0.20	1.50	0.97	0.96
Education	12.5	2.0	12.0	12.0	10-15	0.10	1.80	0.99	0.97
Income	1500	500	1200	1000	500-2500	0.30	2.50	0.96	0.94
Occupation	3.0	1.5	2.5	2.0	1-5	0.25	1.60	0.97	0.95
Health Status	4.0	1.0	4.0	4.0	3-5	0.10	1.20	0.99	0.98
Stress Level	3.5	1.2	3.0	3.0	2-5	0.20	1.40	0.97	0.96
Life Satisfaction	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Resilience	3.0	1.0	2.5	2.5	1-5	0.25	1.50	0.96	0.94
Optimism	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Gratitude	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Forgiveness	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Empathy	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Compassion	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Kindness	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Generosity	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Patience	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Self-control	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Emotional Stability	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Psychological Well-being	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Life Purpose	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Meaning in Life	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Existential Well-being	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Transcendental Well-being	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Humanistic Well-being	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Existential Well-being	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98
Transcendental Well-being	4.5	1.0	4.0	4.0	3-5	0.15	1.30	0.98	0.97
Humanistic Well-being	4.0	1.0	3.5	3.5	3-5	0.10	1.20	0.99	0.98

Michael J. Heller
Eugene Tu
Glen A. Evans
Ronald G. Sosnowski

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storage
5.0)
5.1)
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44

(A) ADDRESSEE: Lyon & Lyon
(B) STREET: 633 West Fifth Street
(C) CITY: Los Angeles
(D) STATE: California
(E) COUNTRY: USA
(F) ZIP: 90071

(A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

(B) COMPUTER: IBM Compatible

(C) OPERATING SYSTEM: IBM P.C. DOS (Version

(D) SOFTWARE: WordPerfect (Version

(A) APPLICATION NUMBER: 08/271,882
(B) FILING DATE: July 7, 1994
(C) CLASSIFICATION:

(A) APPLICATION NUMBER: 08/146,504

(B) FILING DATE: November 1, 1993

(A) NAME: Murphy, David B.
(B) REGISTRATION NUMBER: 31,125

(C) REFERENCE/DOCKET NUMBER: 207/263

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (213) 489-1600
(B) TELEFAX: (213) 955-0440
(C) TELEX: 67-3510

(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 24
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

GCT AGC CCC TGC TCA TGA GTC TCU 24

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 24
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

AAA AAA AAA AAA AAA AAU 24

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 34
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

CTA CGT GGA CCT GGA GAG GAA GGA GAC TGC CTG U 34

207007-47027650

Variable	Mean	SD	Min	Max
Age	34.5	10.2	22	55
Gender	0.5	0.5	0	1
Marital status	0.6	0.5	0	1
Education	12.5	1.5	10	15
Income	15.2	5.8	10	25
Health status	1.2	0.8	0	2
Stress level	2.5	1.2	1	4
Life satisfaction	3.8	1.5	2	5
Work satisfaction	3.2	1.2	2	4
Family satisfaction	3.5	1.3	2	4
Community satisfaction	3.0	1.1	2	4
Overall satisfaction	3.3	1.2	2	4
Life expectancy	75.5	5.2	65	85
Quality of life	72.5	4.8	60	80
Healthcare access	85.5	3.5	75	95
Healthcare quality	82.5	3.2	70	90
Healthcare cost	78.5	3.8	65	85
Healthcare coverage	88.5	2.5	80	95
Healthcare satisfaction	85.5	3.5	75	95
Healthcare equity	82.5	3.2	70	90
Healthcare efficiency	80.5	3.0	65	85
Healthcare innovation	78.5	3.8	65	85
Healthcare sustainability	75.5	4.2	60	80
Healthcare transparency	72.5	4.5	55	75
Healthcare accountability	70.5	4.0	50	70
Healthcare integrity	68.5	3.5	45	65
Healthcare honesty	65.5	3.0	40	60
Healthcare fairness	62.5	2.5	35	55
Healthcare justice	60.5	2.0	30	50
Healthcare freedom	58.5	1.8	25	45
Healthcare security	55.5	1.5	20	40
Healthcare stability	52.5	1.2	15	35
Healthcare safety	50.5	1.0	10	30
Healthcare peace	48.5	0.8	5	25
Healthcare order	45.5	0.5	0	20
Healthcare harmony	42.5	0.2	0	15
Healthcare unity	40.5	0.1	0	10
Healthcare solidarity	38.5	0.1	0	5
Healthcare cooperation	35.5	0.1	0	5
Healthcare assistance	32.5	0.1	0	5
Healthcare support	30.5	0.1	0	5
Healthcare aid	28.5	0.1	0	5
Healthcare help	25.5	0.1	0	5
Healthcare relief	22.5	0.1	0	5
Healthcare comfort	20.5	0.1	0	5
Healthcare ease	18.5	0.1	0	5
Healthcare convenience	15.5	0.1	0	5
Healthcare simplicity	12.5	0.1	0	5
Healthcare clarity	10.5	0.1	0	5
Healthcare openness	8.5	0.1	0	5
Healthcare honesty	6.5	0.1	0	5
Healthcare integrity	4.5	0.1	0	5
Healthcare honesty	2.5	0.1	0	5
Healthcare honesty	0.5	0.1	0	5

acid

```
(A) LENGTH:                20
(B) TYPE:                   nucleic
                             single
(C) STRANDEDNESS:          single
(D) TOPOLOGY:              linear
```

GAG TTC AGC AAA TTT GGA GU

20

acid

```
(A) LENGTH:                20
(B) TYPE:                   nucleic
                               acid
(C) STRANDEDNESS:          single
(D) TOPOLOGY:              linear
```

20
nucleic

single
linear

CGT AGA ACT CCT CAT CTC CU

20

acid

```
(A) LENGTH: 18
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear
```

18
nucleic

single
linear

GTC TCC TTC CTC TCC AGU

18

acid

```
(A) LENGTH:                20
(B) TYPE:                   nucleic

(C) STRANDEDNESS:          single
(D) TOPOLOGY:              linear
```

20
nucleic

single
linear

20

(i) SEQUENCE CHARACTERISTICS:

acid

18

(i) SEQUENCE CHARACTERISTICS:

acid

22

(i) SEQUENCE CHARACTERISTICS:

acid

20

(2) INFORMATION FOR SEQ ID NO: 11:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 20
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

TGT TTG CCT GTT CTC AGA CU 20

(2) INFORMATION FOR SEQ ID NO: 12:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 20
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

CAT CGC TGT GAC AAA ACA TU 20

(2) INFORMATION FOR SEQ ID NO: 13:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 20
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

TGC GAG CTG CAG TCA GAC AT 20

(2) INFORMATION FOR SEQ ID NO: 14:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 18
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

130100747021550

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

GAG AGA CTC ATG AGC AGG

18

(2) INFORMATION FOR SEQ ID NO: 15:

(i) SEQUENCE CHARACTERISTICS:

acid	(A) LENGTH:	18
	(B) TYPE:	nucleic
	(C) STRANDEDNESS:	single
	(D) TOPOLOGY:	linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

CCT GCT CAT GAG TCT CTC

18

(2) INFORMATION FOR SEQ ID NO: 16:

(i) SEQUENCE CHARACTERISTICS:

acid	(A) LENGTH:	19
	(B) TYPE:	nucleic
	(C) STRANDEDNESS:	single
	(D) TOPOLOGY:	linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

TTT TTT TTT TTT TTT TTT T

19

(2) INFORMATION FOR SEQ ID NO: 17:

(i) SEQUENCE CHARACTERISTICS:

acid	(A) LENGTH:	33
	(B) TYPE:	nucleic
	(C) STRANDEDNESS:	single
	(D) TOPOLOGY:	linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

CAG GCA GTC TCC TTC CTC TCC AGG TCC ACG TAG

33

(2) INFORMATION FOR SEQ ID NO: 18:

(i) SEQUENCE CHARACTERISTICS:

TCF001"4T02T650

acid (A) LENGTH: 19
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 18:
CTC CAA ATT TGC TGA ACT C 19

(2) INFORMATION FOR SEQ ID NO: 19:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 19
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 19:
GGA GAT GAG GAG TTC TAC G 19

(2) INFORMATION FOR SEQ ID NO: 20:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 17
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 20:
CTG GAG AGG AAG GAG AC 17

(2) INFORMATION FOR SEQ ID NO: 21:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 19
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

19

(i) SEQUENCE CHARACTERISTICS:

```

acid      (A) LENGTH:                17
          (B) TYPE:                 nucleic
          (C) STRANDEDNESS:         single
          (D) TOPOLOGY:             linear

```

17

(i) SEQUENCE CHARACTERISTICS:

```

      (A) LENGTH:                21
      (B) TYPE:                  nucleic
acid
      (C) STRANDEDNESS:          single
      (D) TOPOLOGY:              linear

```

21

(i) SEQUENCE CHARACTERISTICS:

```

acid      (A) LENGTH:                19
          (B) TYPE:                 nucleic
          (C) STRANDEDNESS:         single
          (D) TOPOLOGY:             linear

```

19

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 19
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 25:

GTC TGA GAA CAG GCA AAC A 19

(2) INFORMATION FOR SEQ ID NO: 26:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 19
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 26:

ATG TTT TGT CAC AGC GAT G 19

(2) INFORMATION FOR SEQ ID NO: 27:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 30
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 27:

GGT GGT GGG CGC CGG CGG TGT GGG CAA GAU 30

(2) INFORMATION FOR SEQ ID NO: 28:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 30
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

0912014-10010

GGT GGT GGG CGC CGT CGG TGT GGG CAA GAU

30

(2) INFORMATION FOR SEQ ID NO: 29:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 15
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 29:

CC GCG GCC GCC ACA C 15

(2) INFORMATION FOR SEQ ID NO: 30:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 15
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 30:

CC GCG GCA GCC ACA C 15

(2) INFORMATION FOR SEQ ID NO: 31:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 15
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 31:

CC GTG GCA GCC ACA C 15

(2) INFORMATION FOR SEQ ID NO: 32:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 30
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 32:

GGT GGT GGG CGC CGG CGG TGT GGG CAA GAU 30

(2) INFORMATION FOR SEQ ID NO: 33:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 29
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

GGT GGT GGG CGC CGG CGG TGT GGG CAA GA 29

(2) INFORMATION FOR SEQ ID NO: 34:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 22
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

TGC CCA CAC CGC CGG CGC CCA C 22

(2) INFORMATION FOR SEQ ID NO: 35:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 22
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 35:

TOPP-4T02T660

22

(i) SEQUENCE CHARACTERISTICS:

```

acid      (A) LENGTH:                22
          (B) TYPE:                  nucleic
          (C) STRANDEDNESS:          single
          (D) TOPOLOGY:              linear

```

TGC CCA CAC CGA CGG TGC CCA C 22

(i) SEQUENCE CHARACTERISTICS:

```

acid      (A) LENGTH:          7
          (B) TYPE:          nucleic
          (C) STRANDEDNESS:   single
          (D) TOPOLOGY:      linear

```

ACA CCG C 7

(i) SEQUENCE CHARACTERISTICS:

```

acid      (A) LENGTH:          7
          (B) TYPE:          nucleic
          (C) STRANDEDNESS:   single
          (D) TOPOLOGY:      linear

```

ACA ACG C 7

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31

acid (B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 39:

CCA GTC ACG ACG TTG TAA AAC GAC GGC CAG U 31

(2) INFORMATION FOR SEQ ID NO: 40:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 31
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 40:

GTA ATC ATG GTC ATA GCT GTT TCC TGT GTG U 31

(2) INFORMATION FOR SEQ ID NO: 41:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 40
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 41:

GCA TGC CTG CAG GTC GAC TCT AGA GGA TCC CCG GGT ACC G 40

(2) INFORMATION FOR SEQ ID NO: 42:

(i) SEQUENCE CHARACTERISTICS:

acid (A) LENGTH: 40
(B) TYPE: nucleic
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 42:

TGC CAA GCT TGG CTG CAG GTC GAC GGA TCC CCG GGA ATT C 40

[illegible]

acid

```
(A) LENGTH:                24
(B) TYPE:                   nucleic
(C) STRANDEDNESS:          single
(D) TOPOLOGY:              linear
```

AAA TTG TTA TCC GCT CAC AAT TGC

24

acid

```
(A) LENGTH:                24
(B) TYPE:                  nucleic
(C) STRANDEDNESS:         single
(D) TOPOLOGY:             linear
```

ACA CAA CAT ACG AGC CGG AAG CAT

24